

Fig. 1A

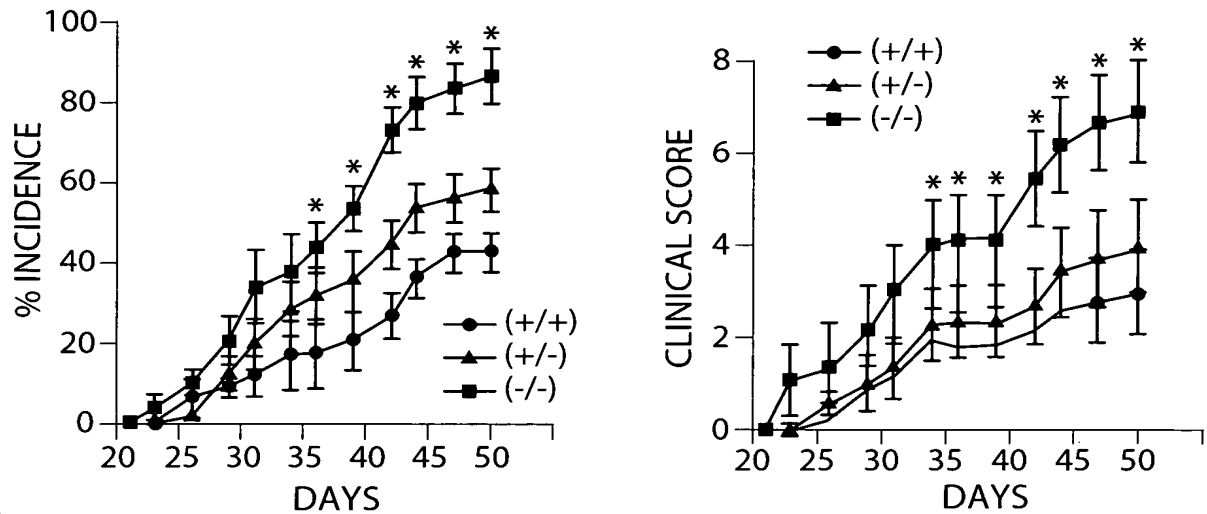
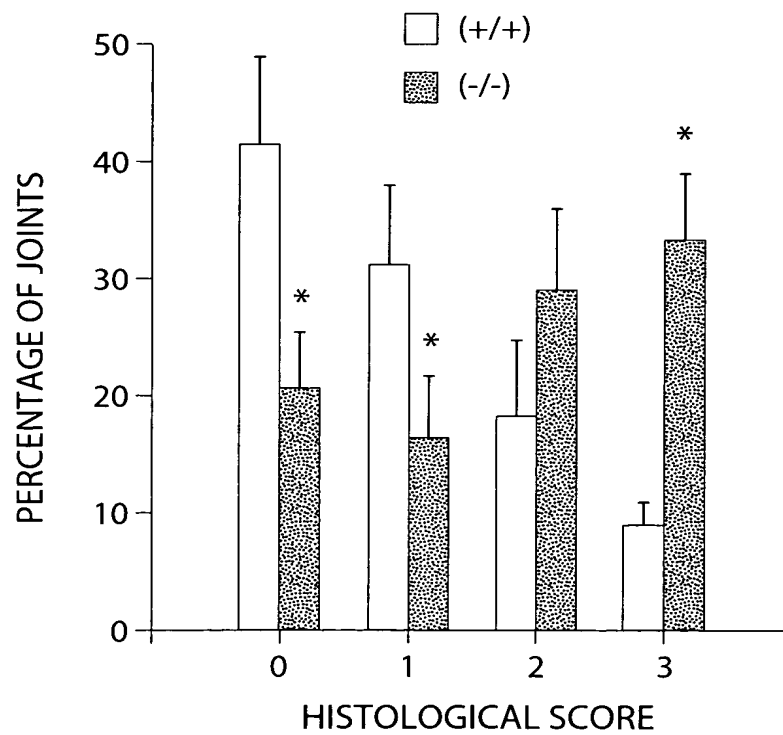
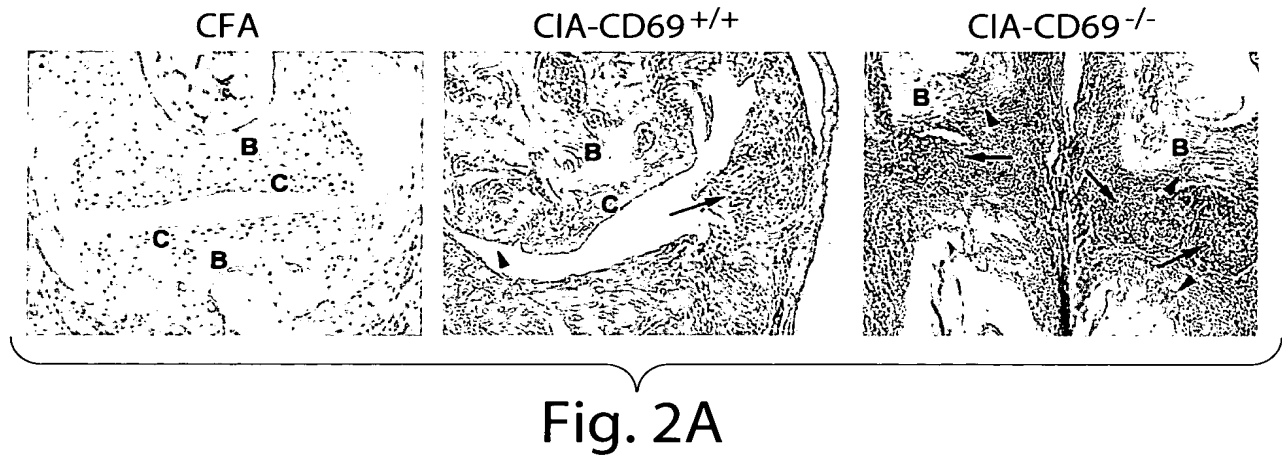


Fig. 1B

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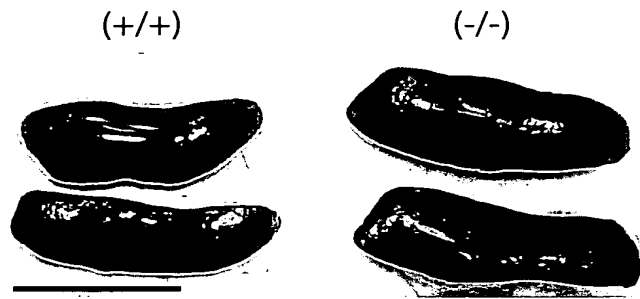


Fig. 3A

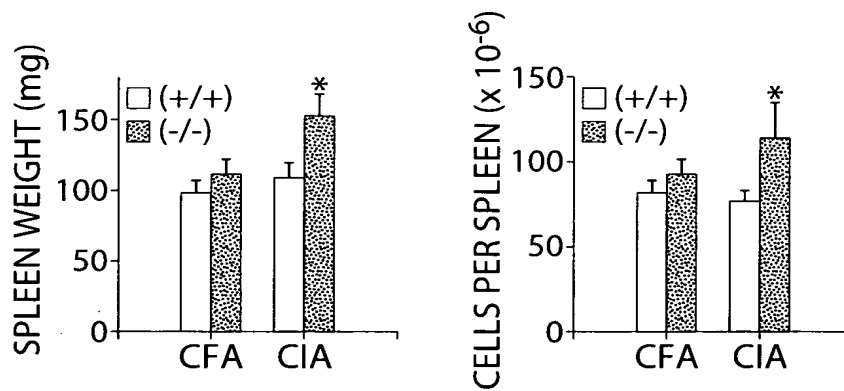


Fig. 3B

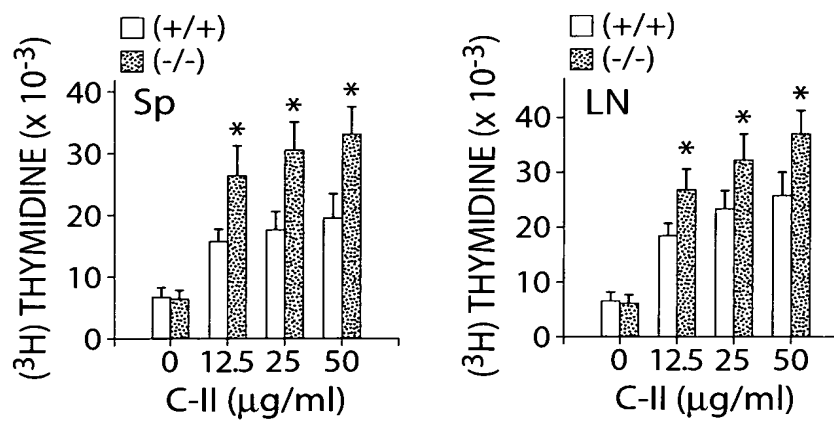


Fig. 3C

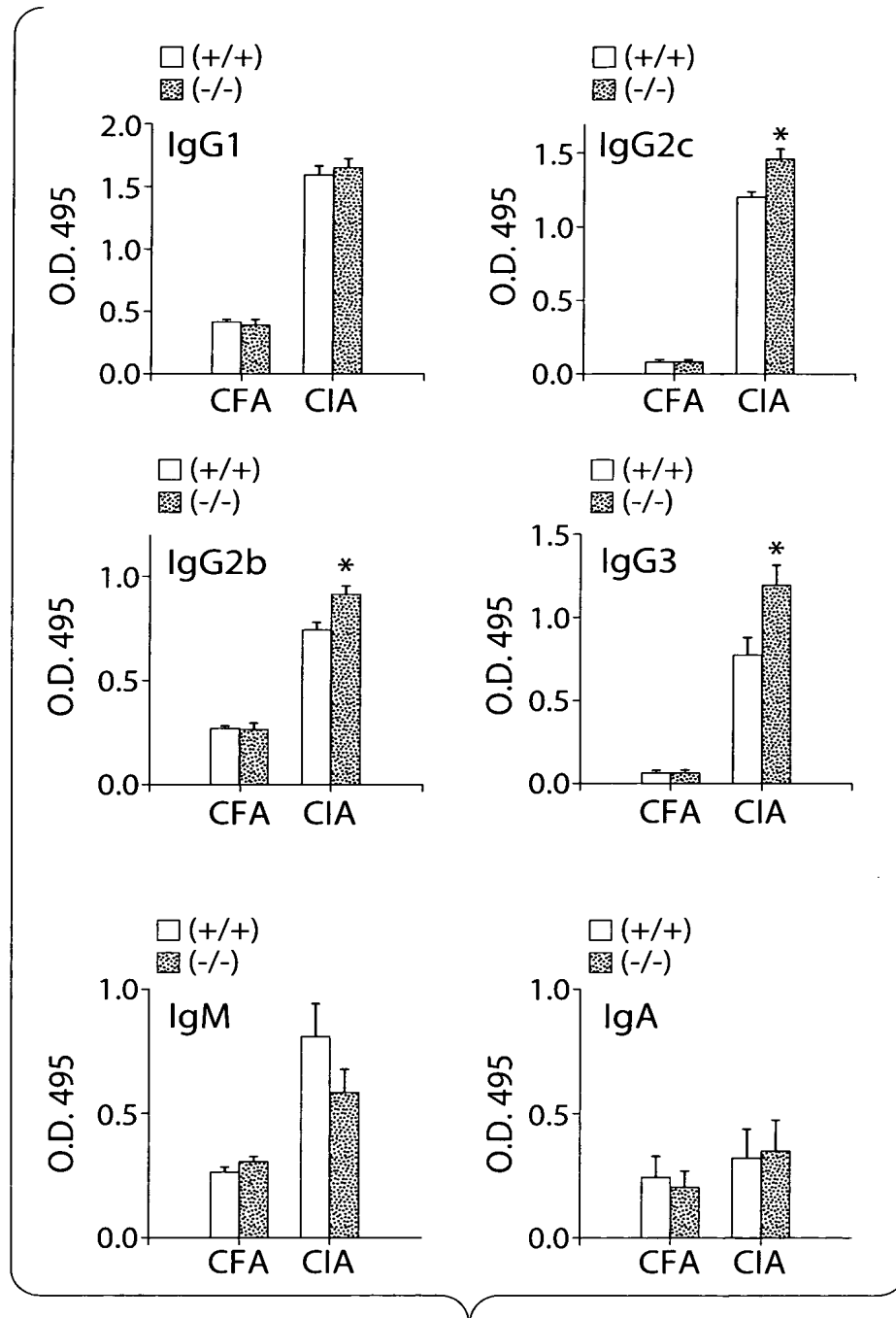


Fig. 3D

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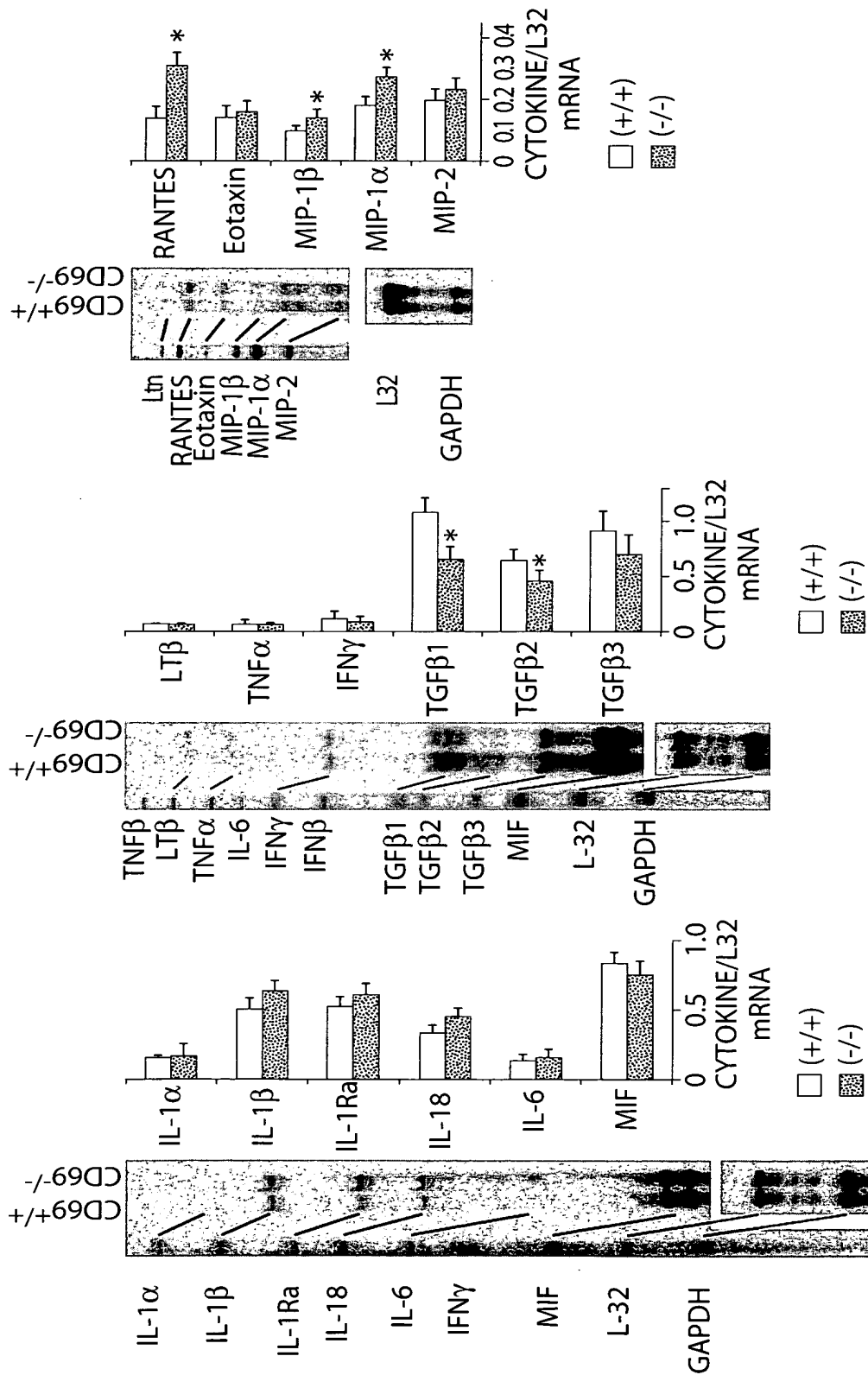


Fig. 4

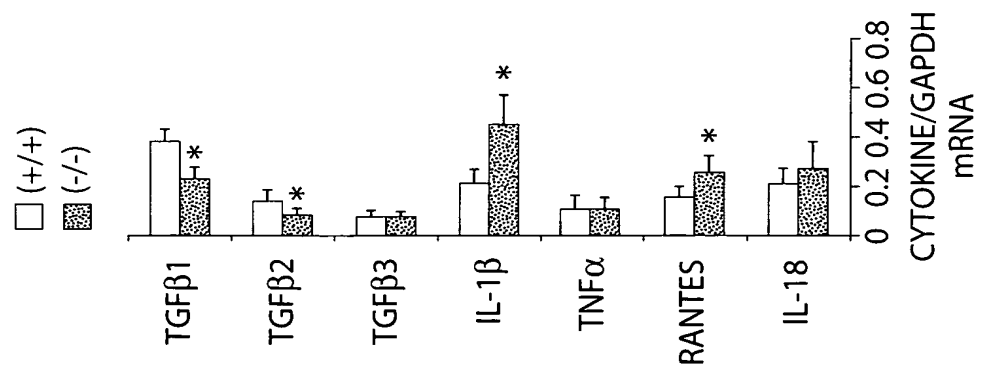


Fig. 5A

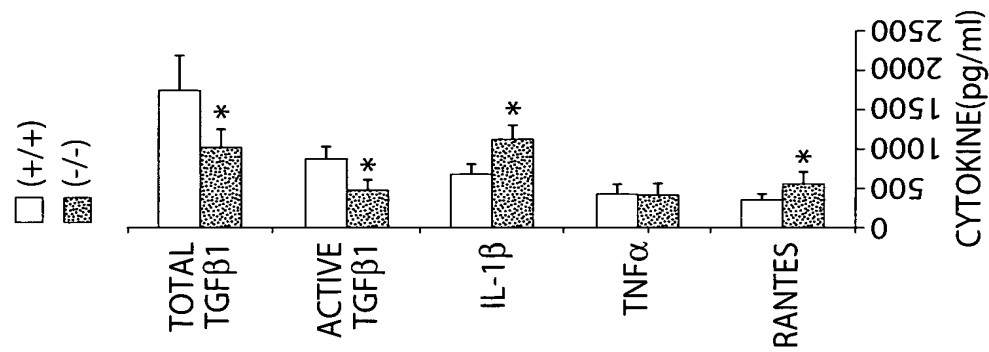


Fig. 5B

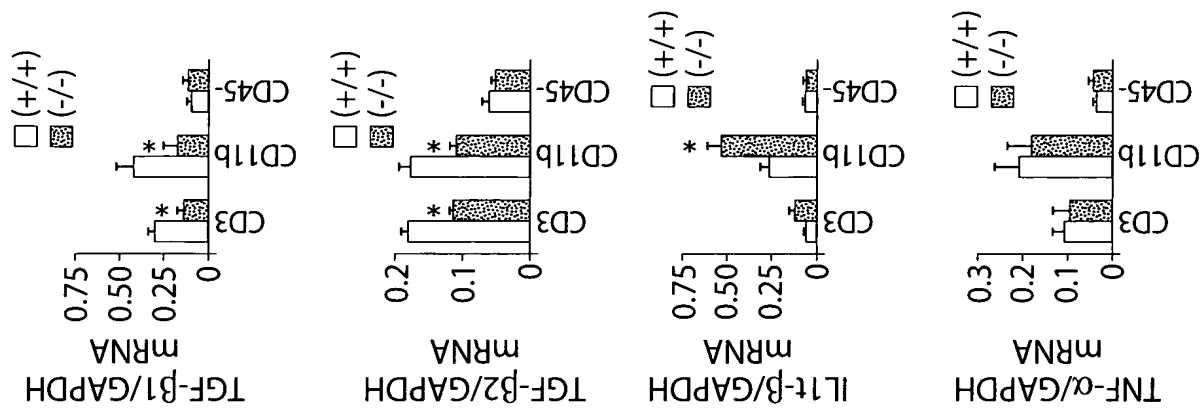


Fig. 5C

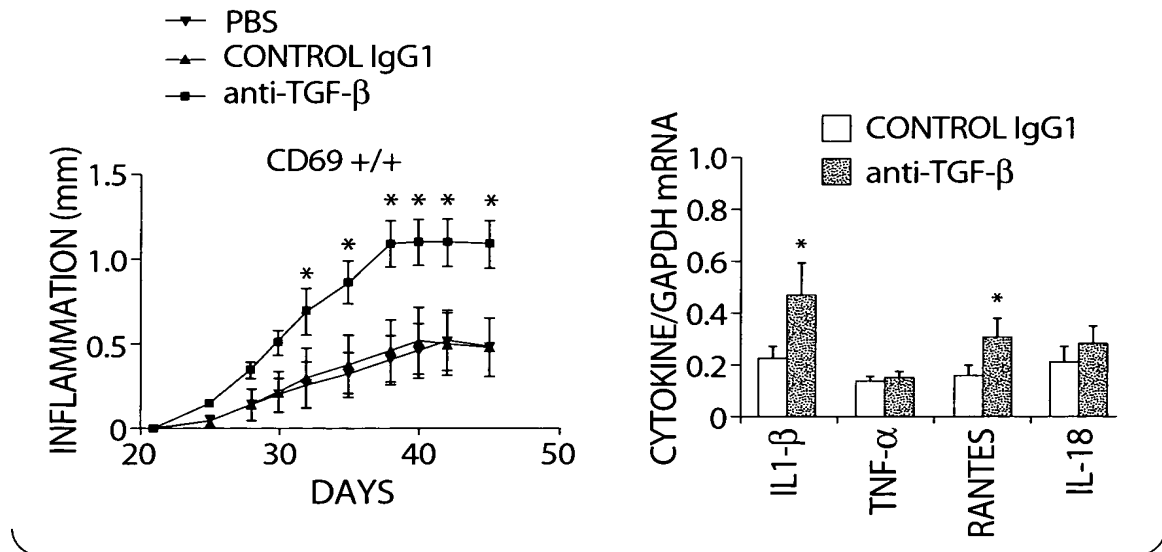


Fig. 6A

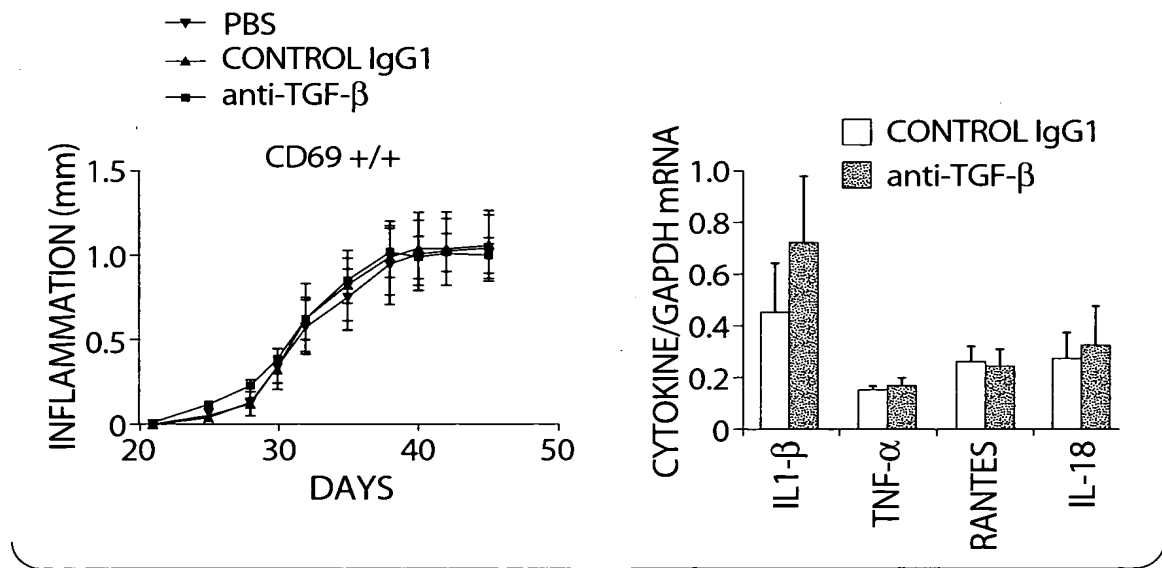


Fig. 6B

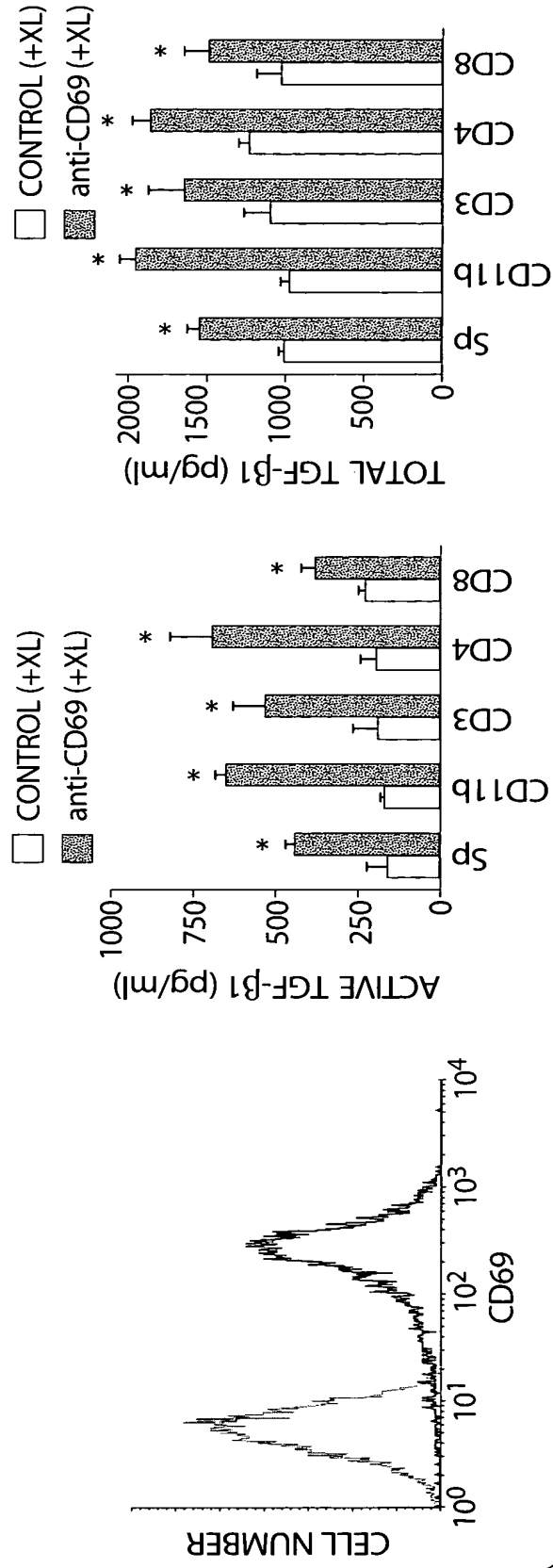


Fig. 7A

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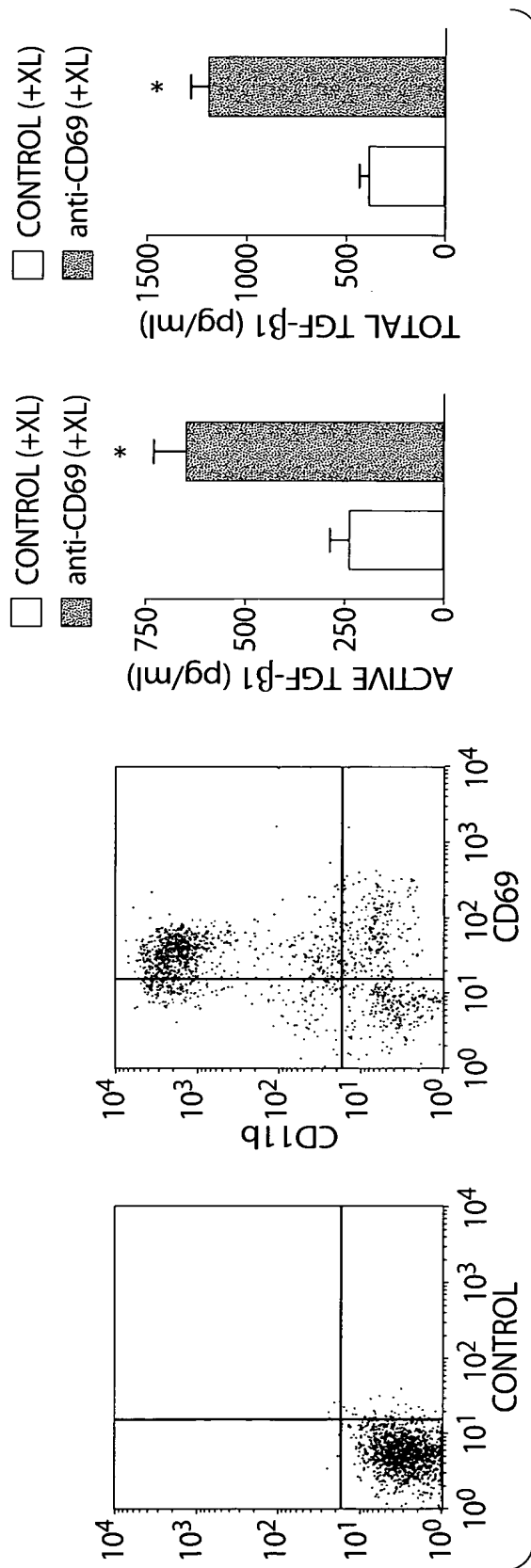


Fig. 7B

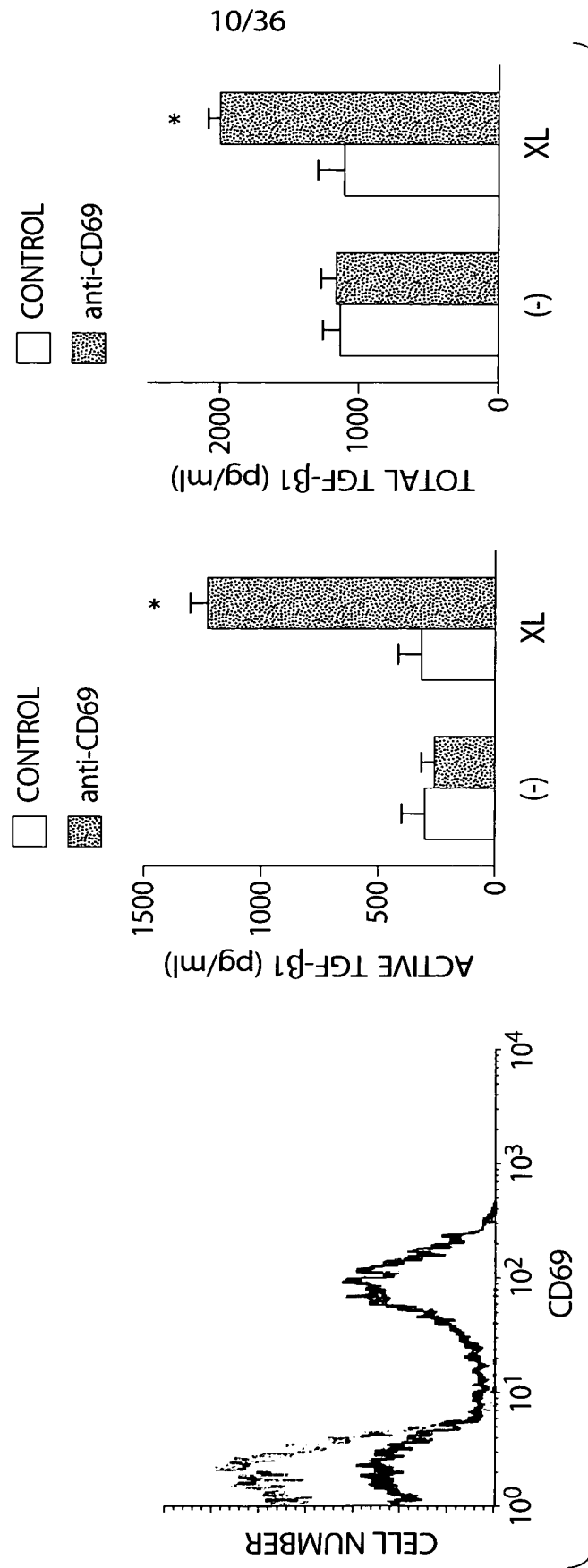
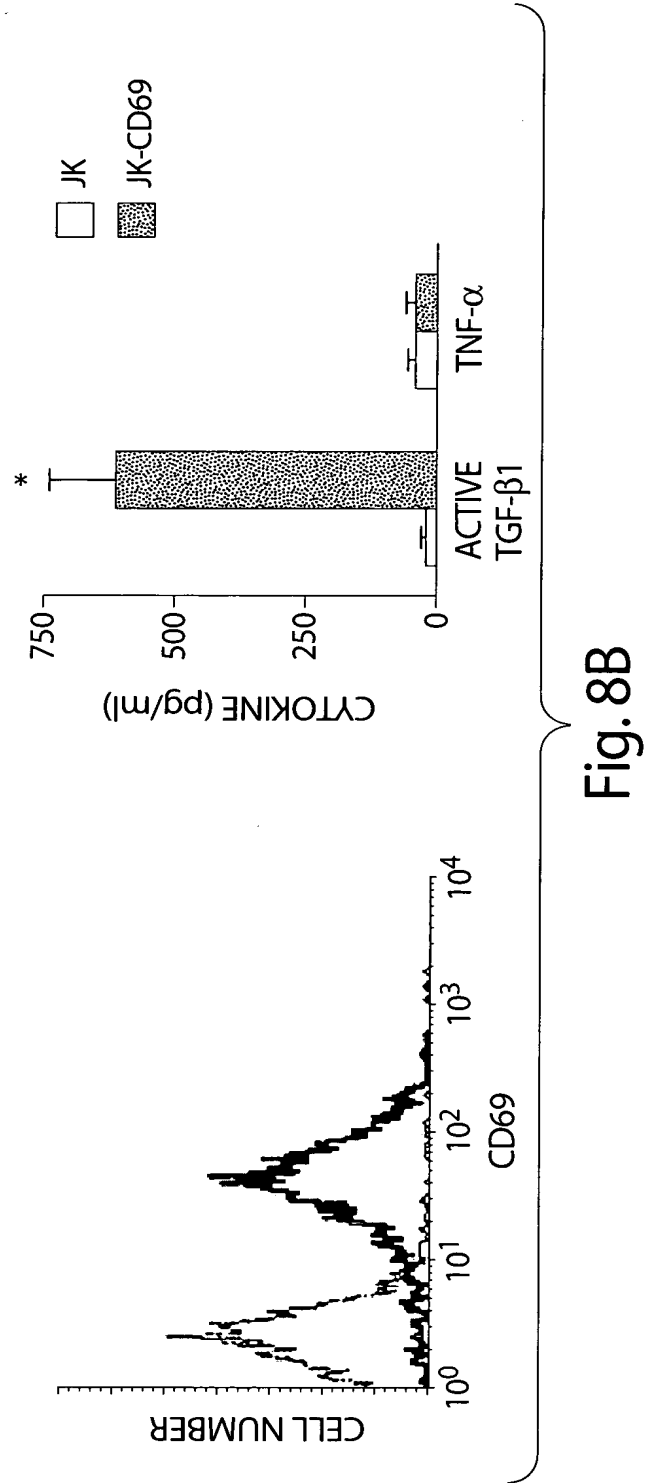


Fig. 8A



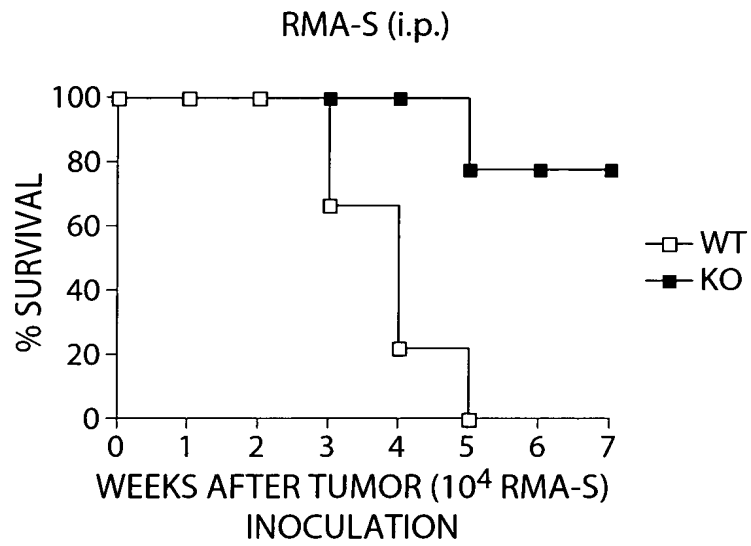


Fig. 9A

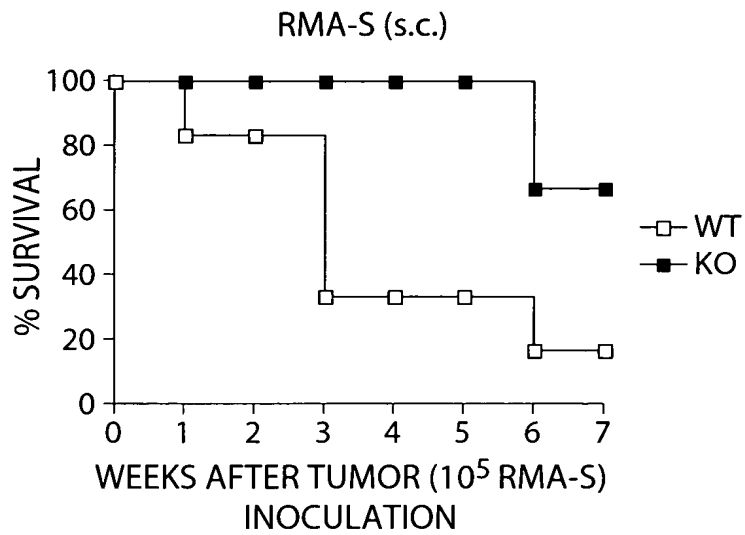


Fig. 9B

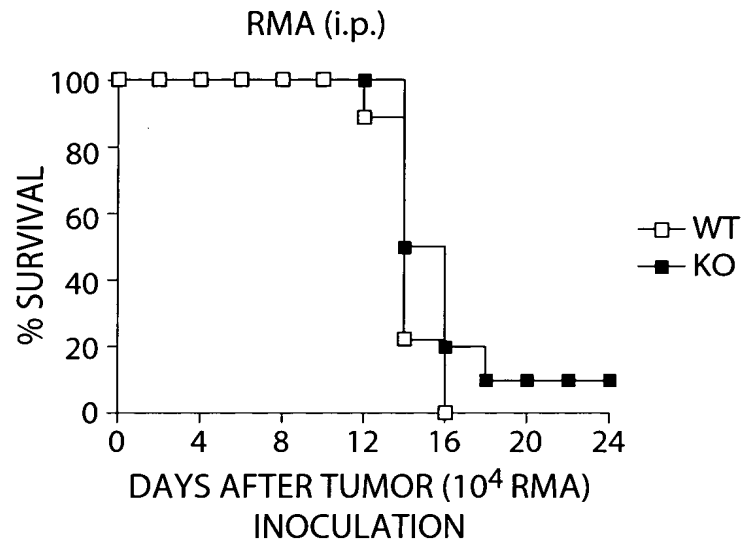


Fig. 9C

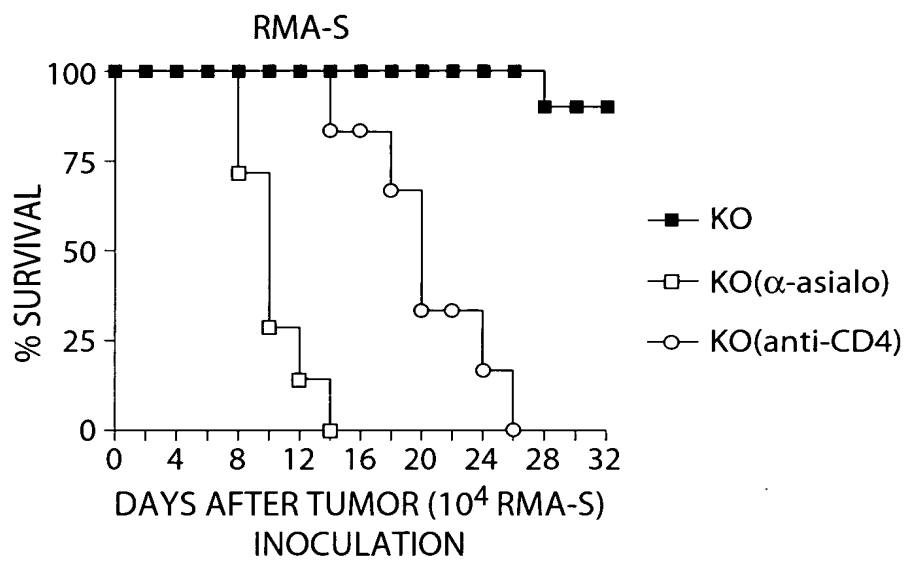


Fig. 9D

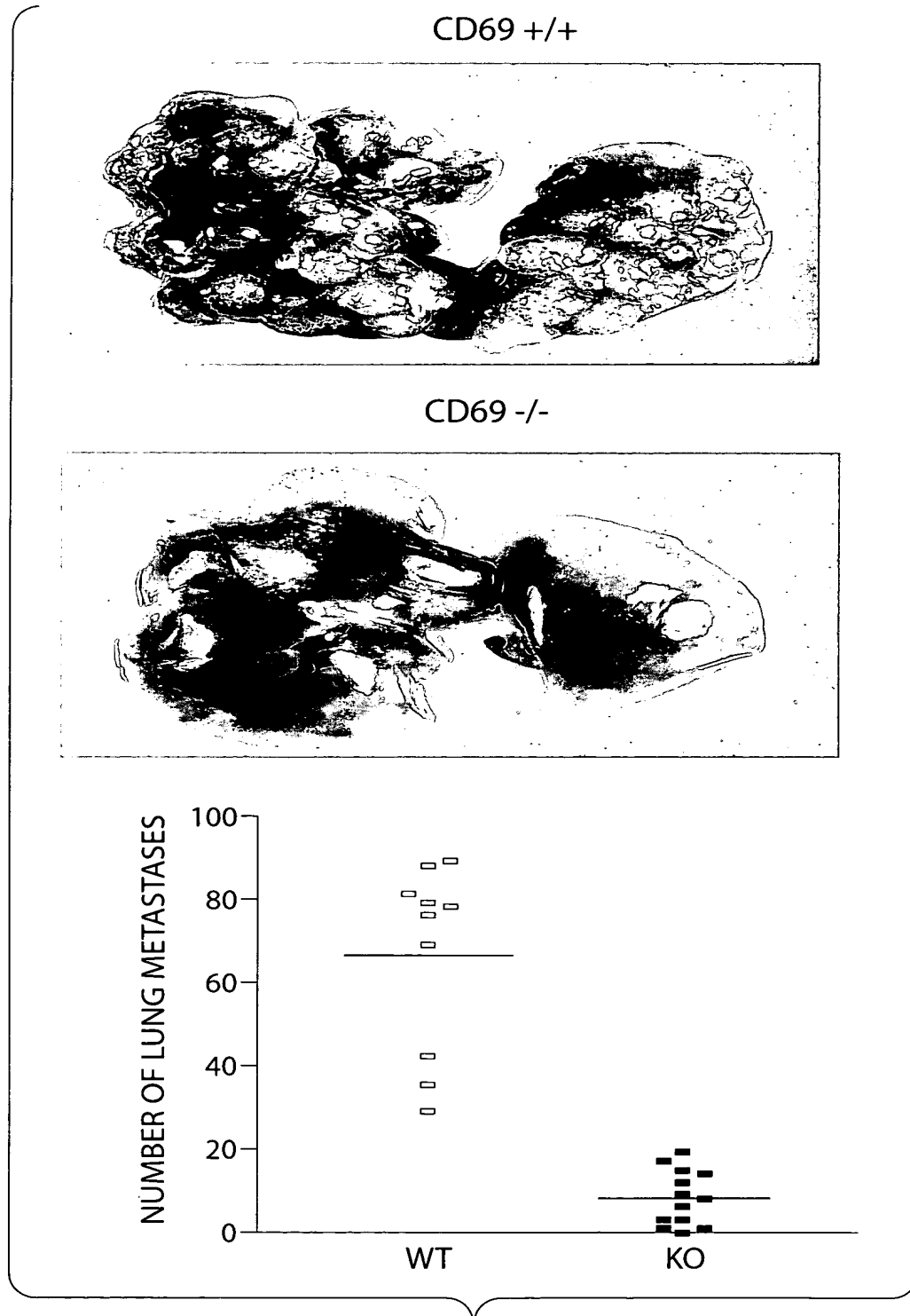


Fig. 9E

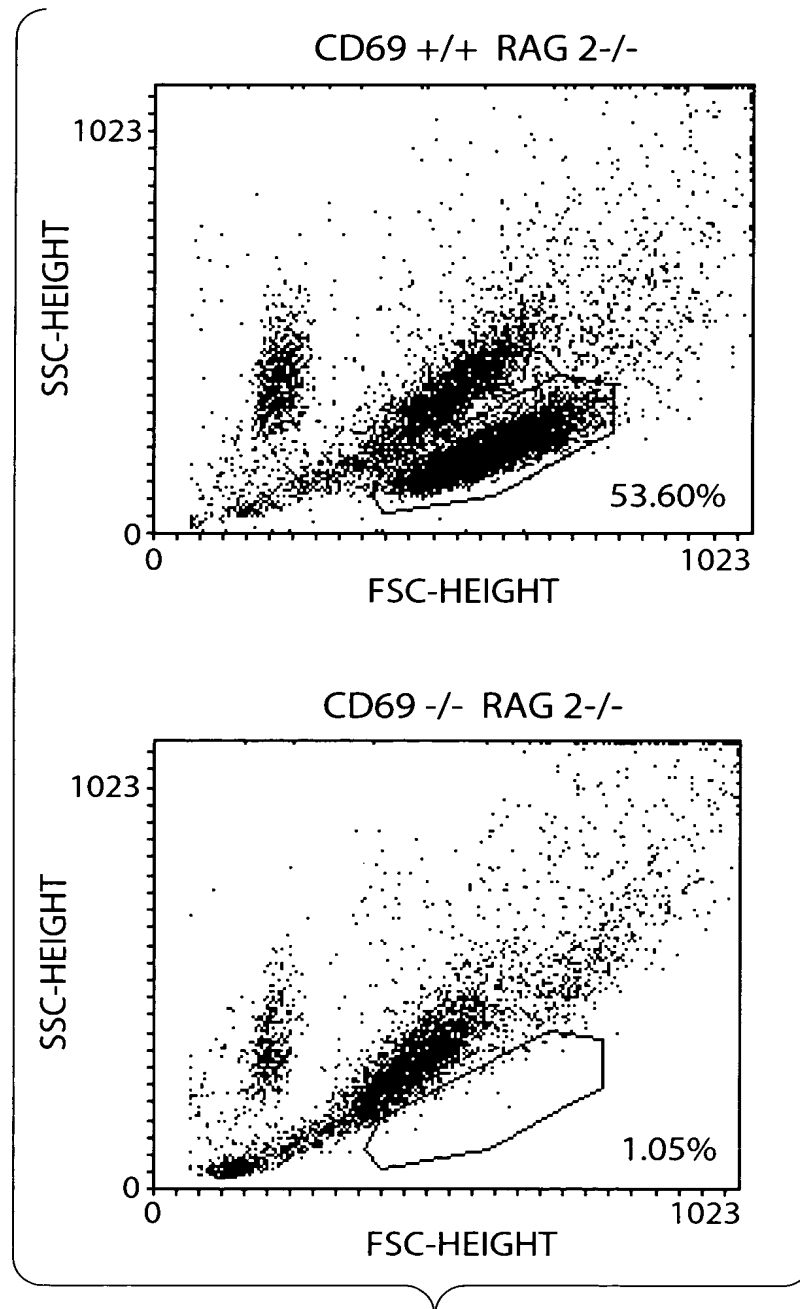


Fig. 10A

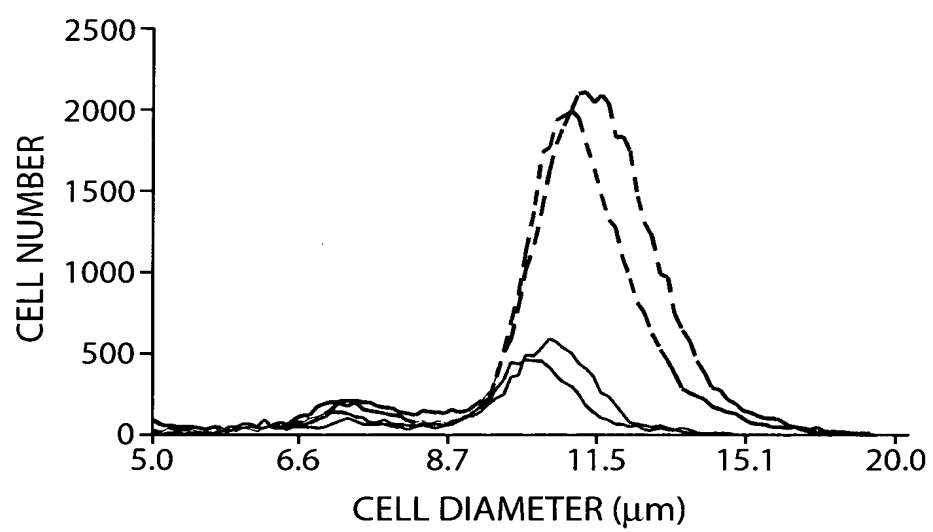


Fig. 10B

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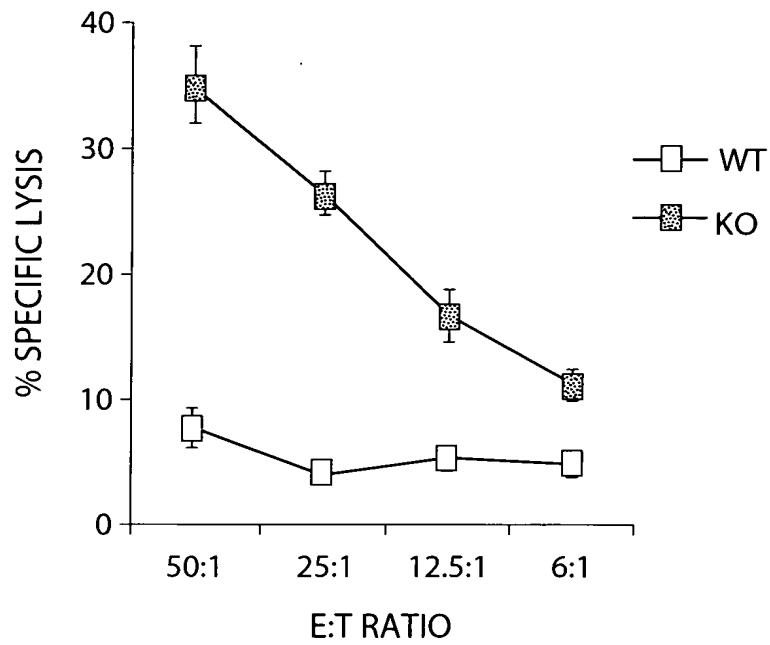


Fig. 11A

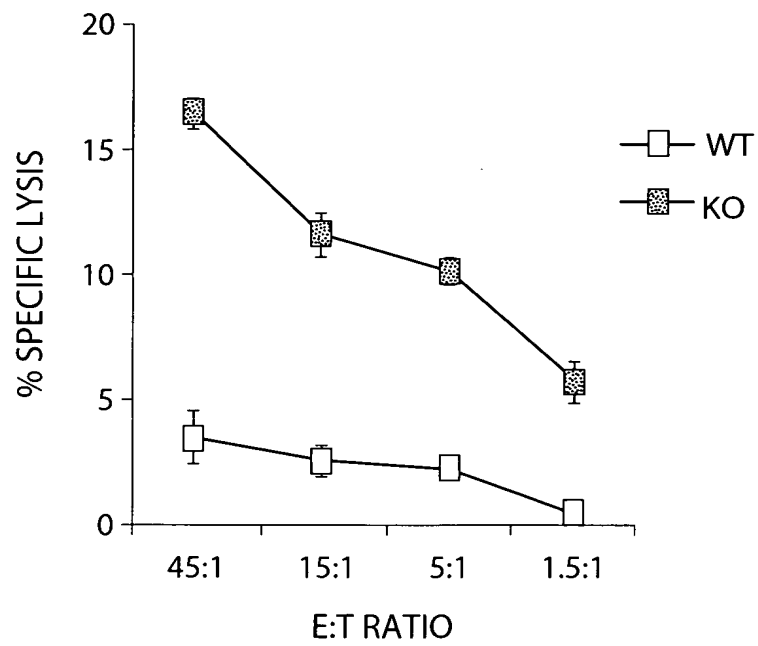


Fig. 11B

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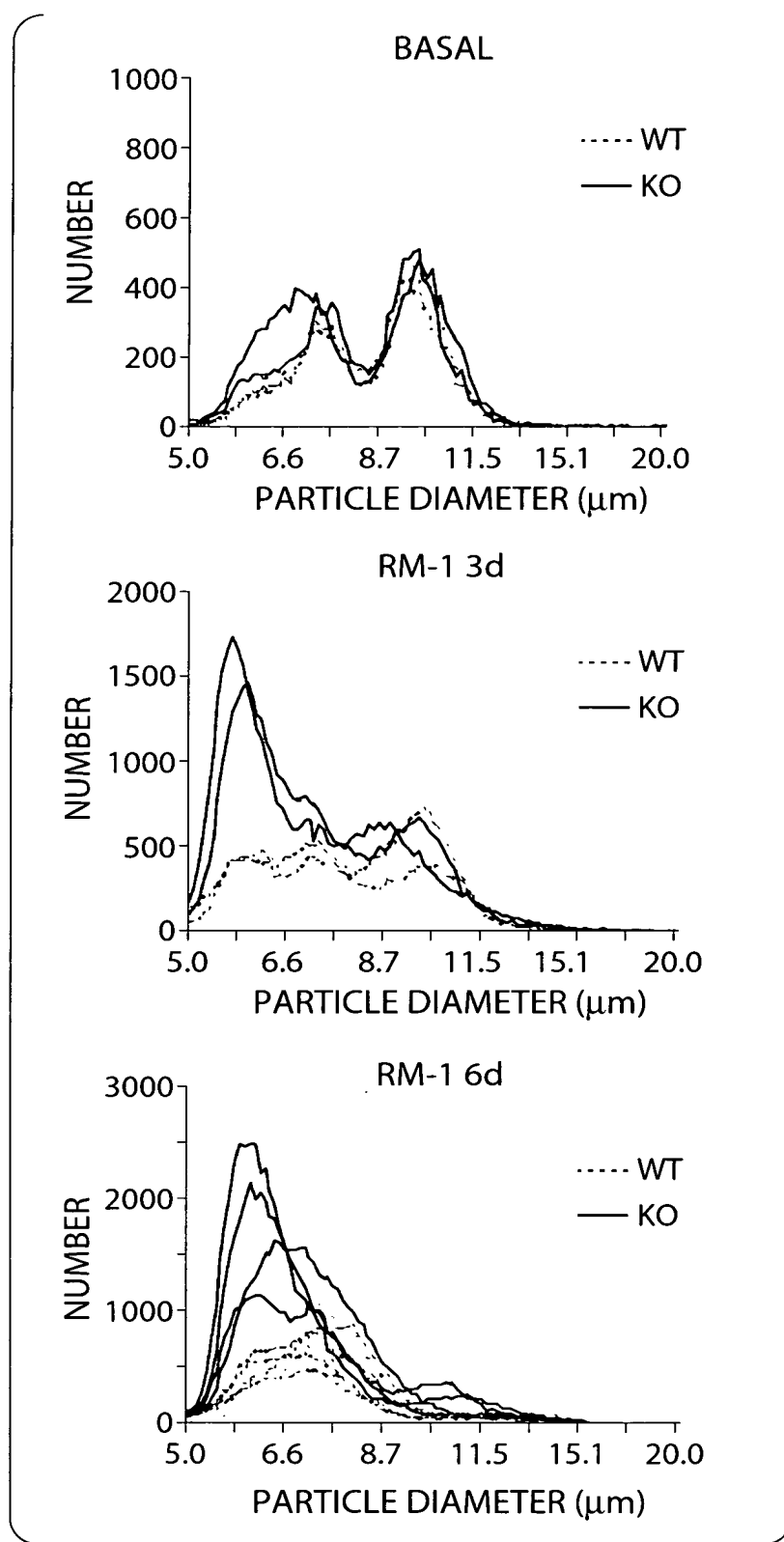


Fig. 11C

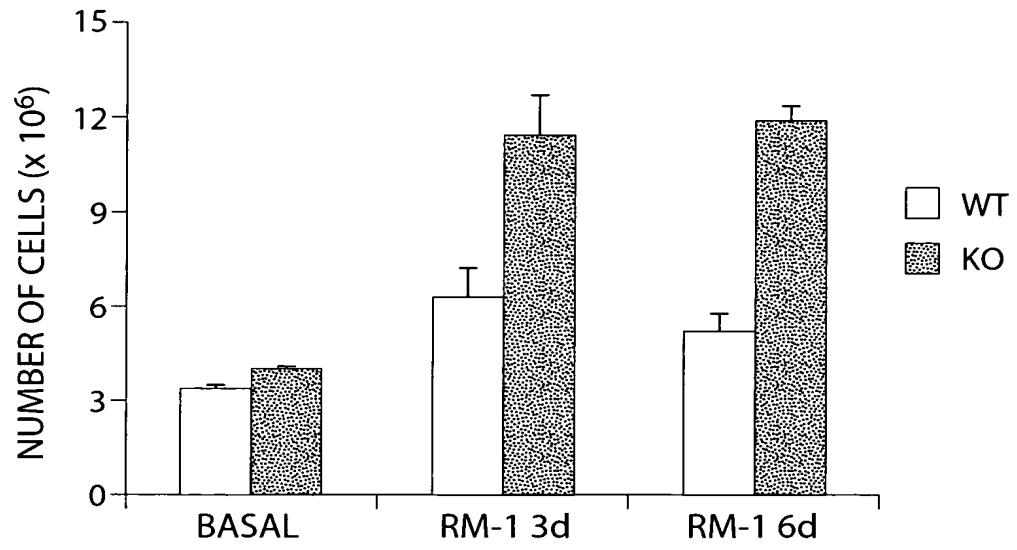


Fig. 11D

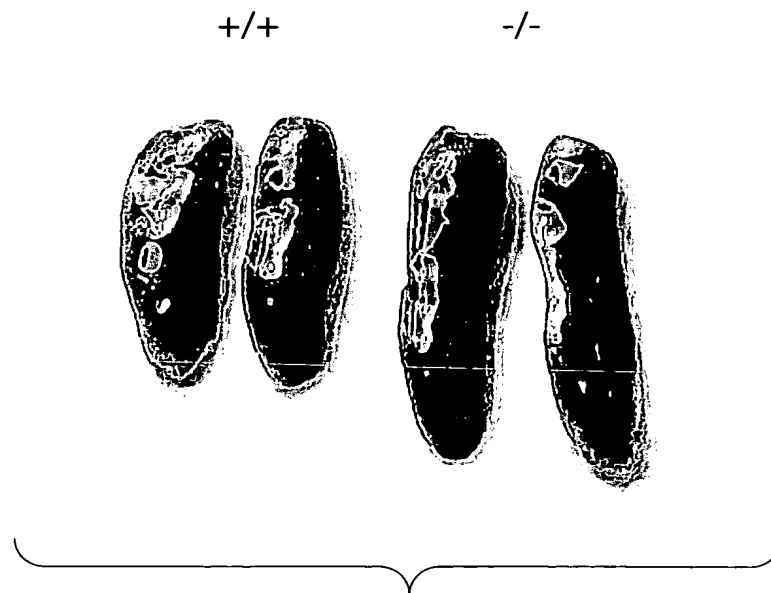


Fig. 11E

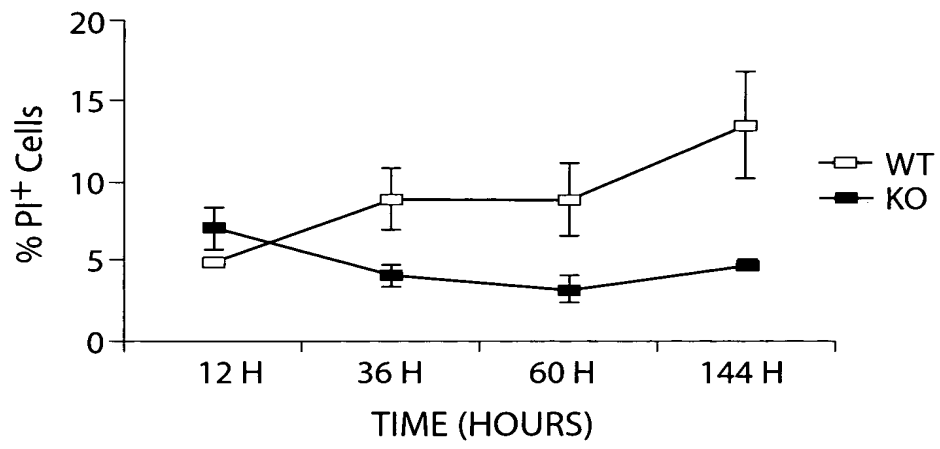


Fig. 12A

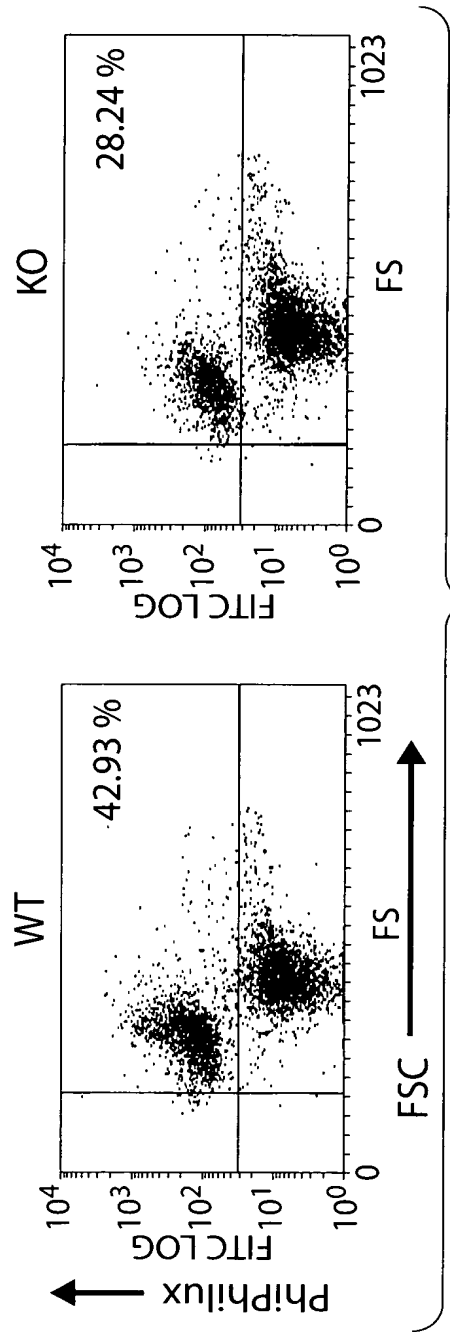


Fig. 12B

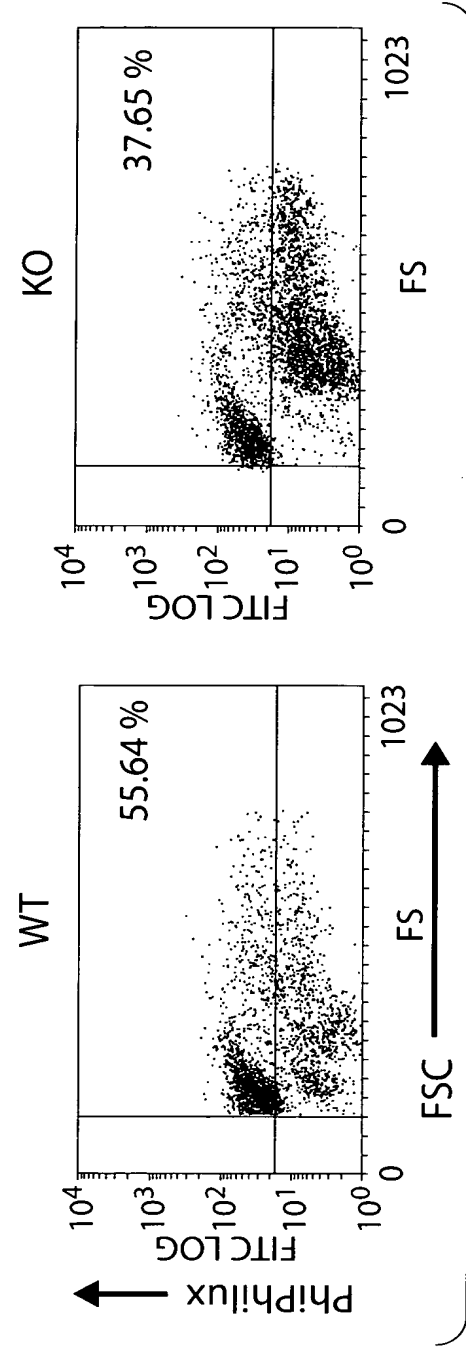
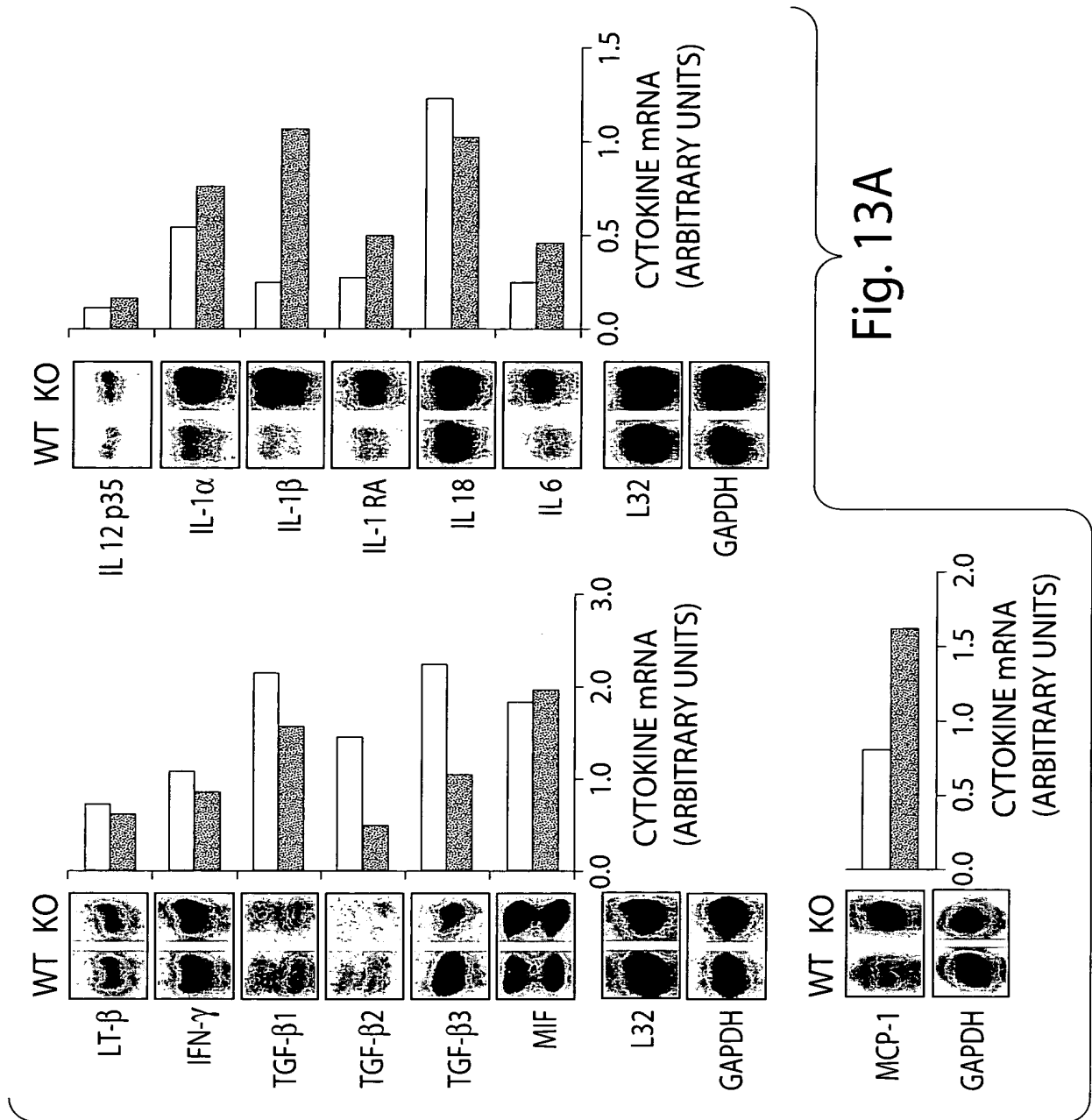


Fig. 12C



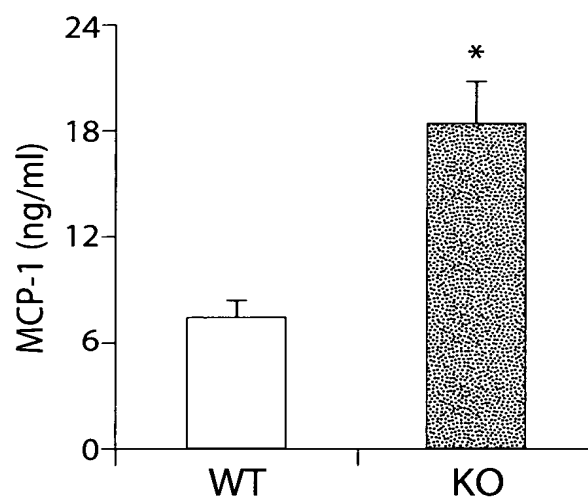


Fig. 13B

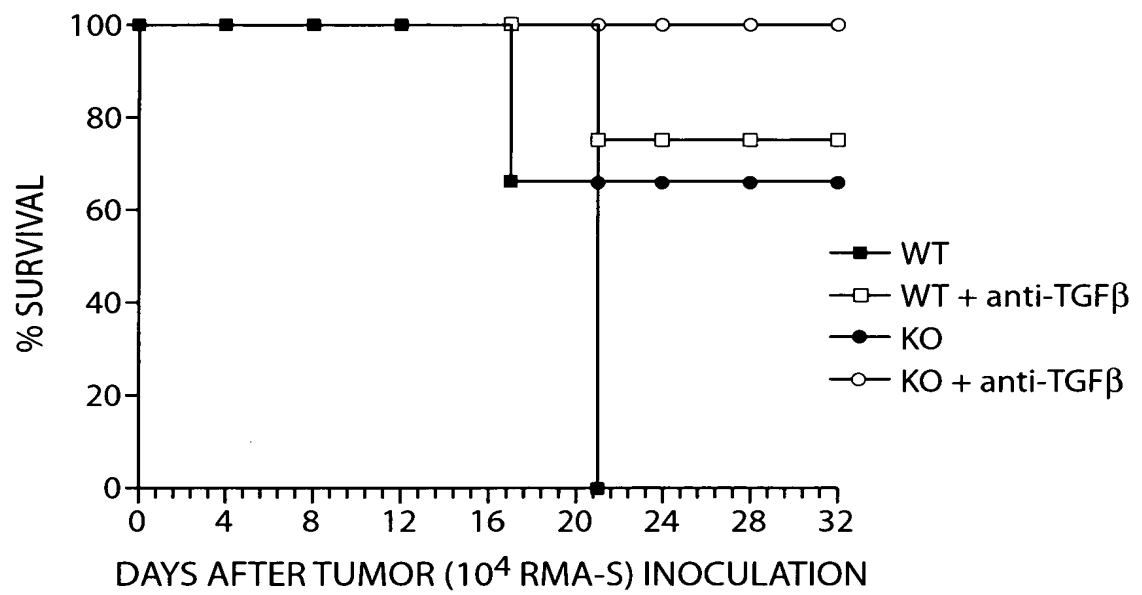


Fig. 14

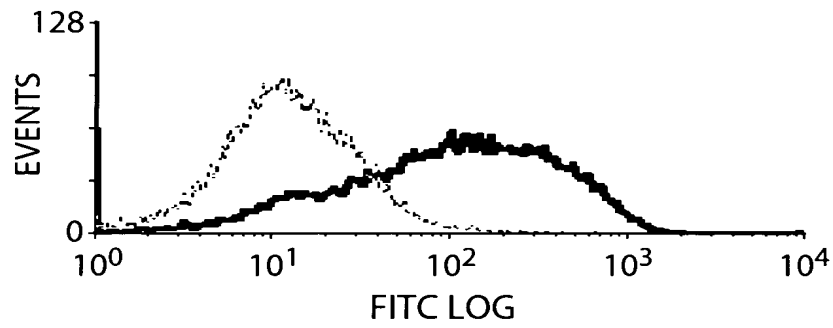


Fig. 15A

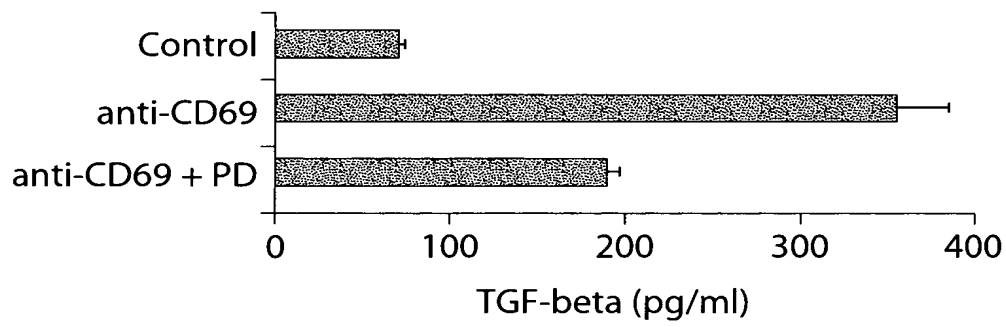


Fig. 15B

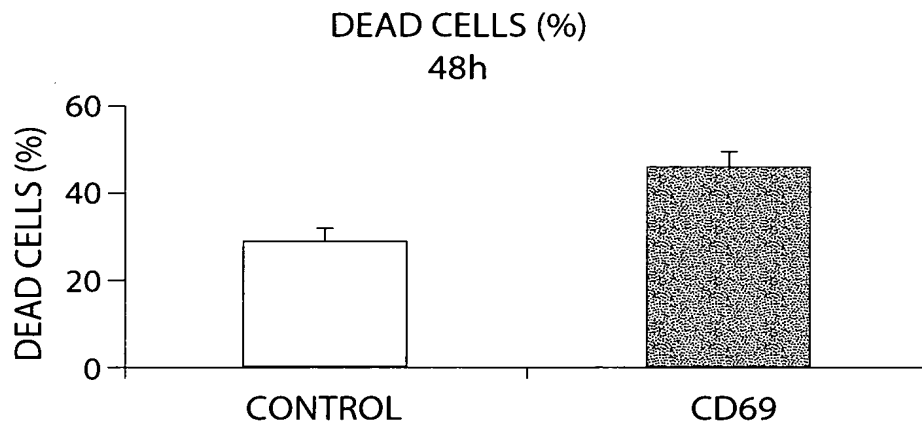


Fig. 15C

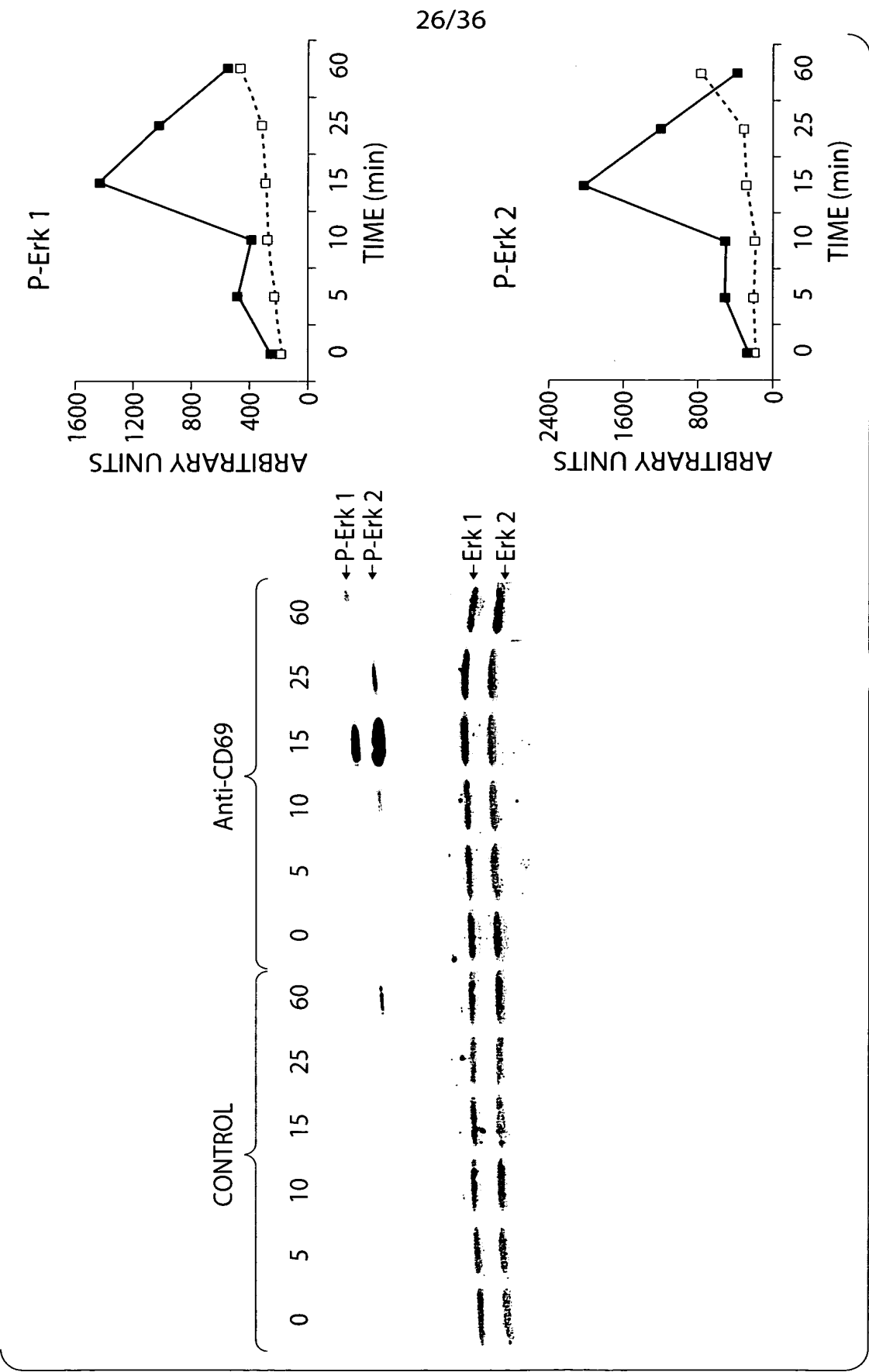


Fig. 15D

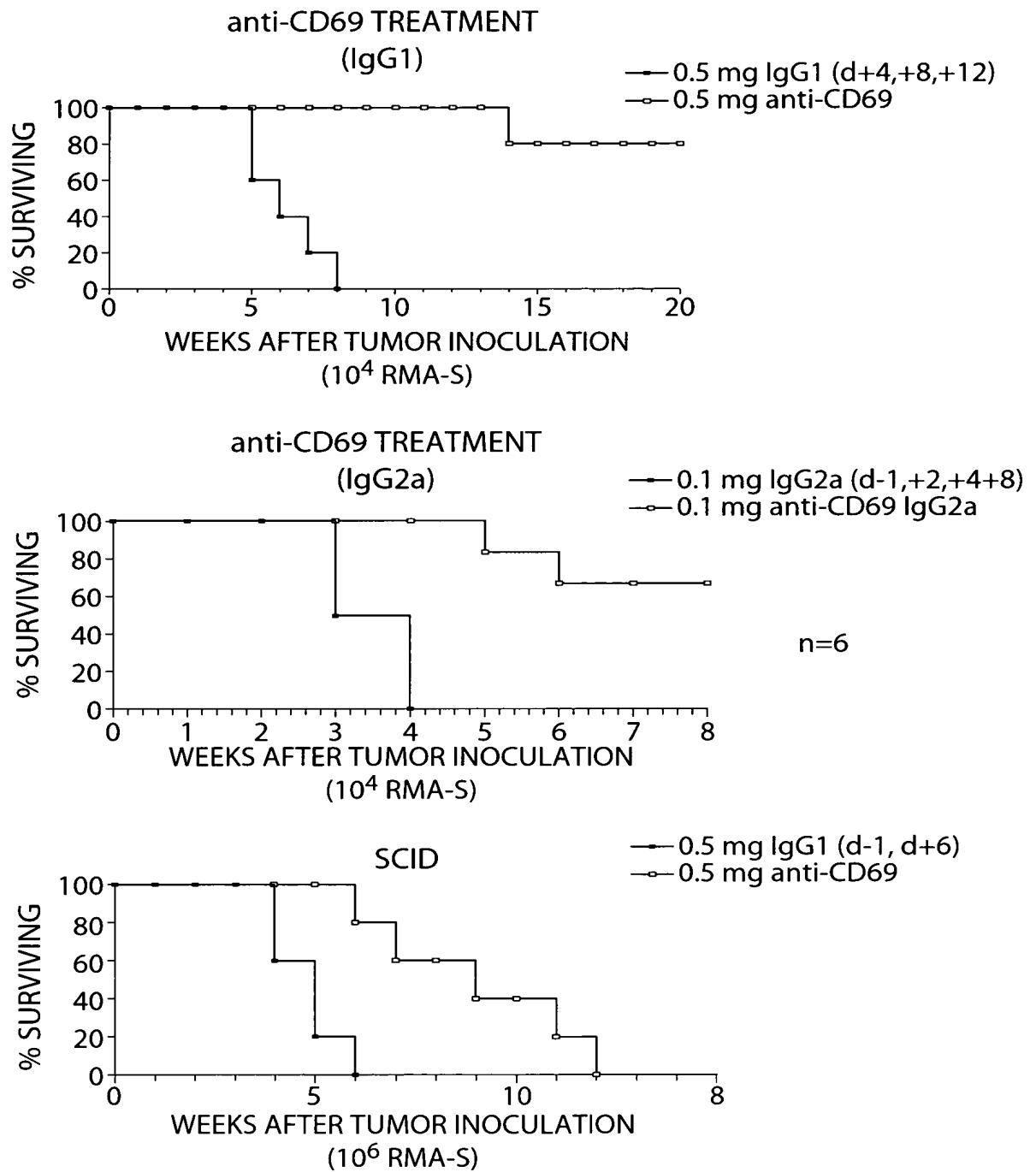


Fig. 16

C57bl/6

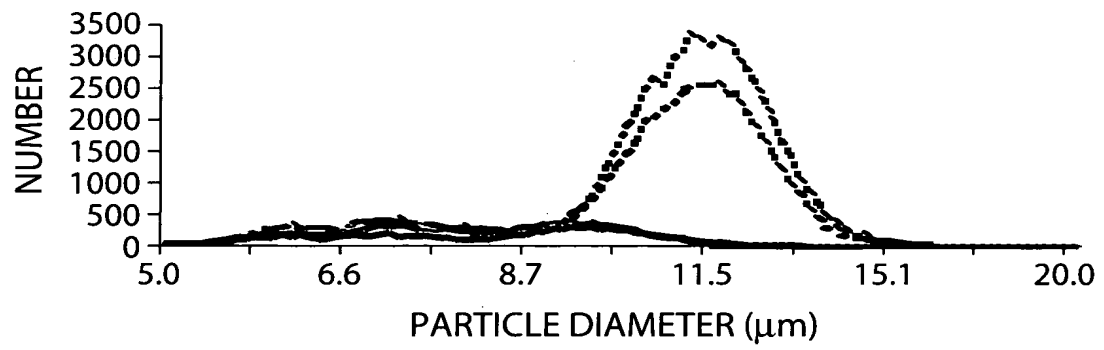


Fig. 17A

Rag1-/-

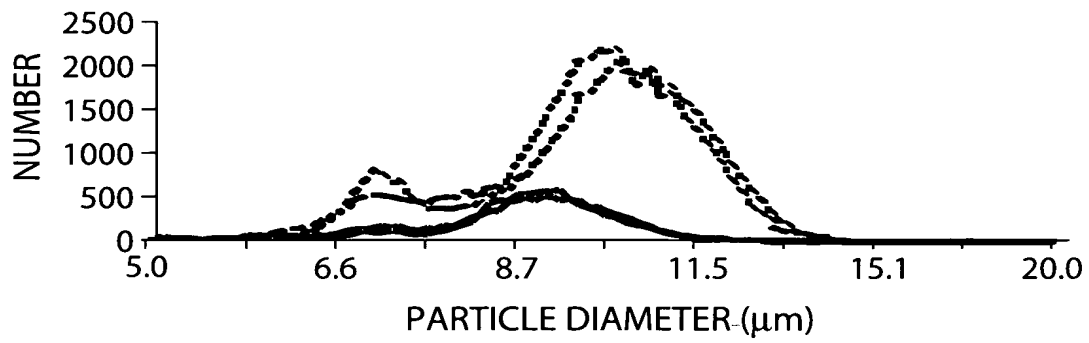


Fig. 17B

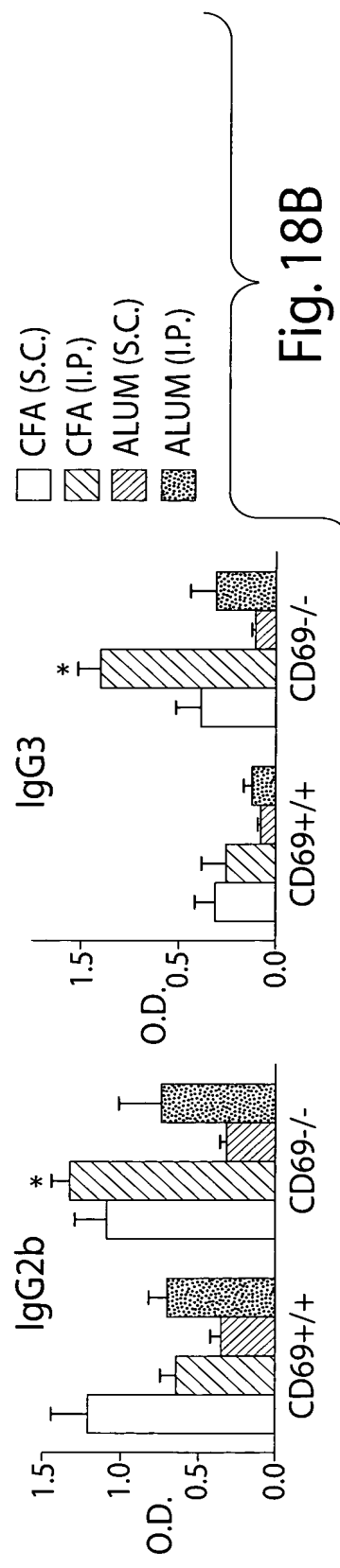
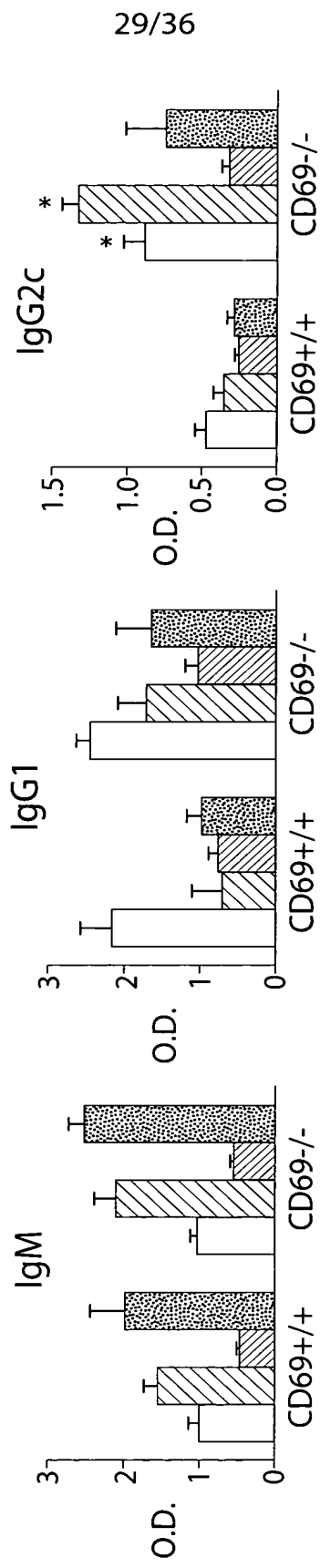
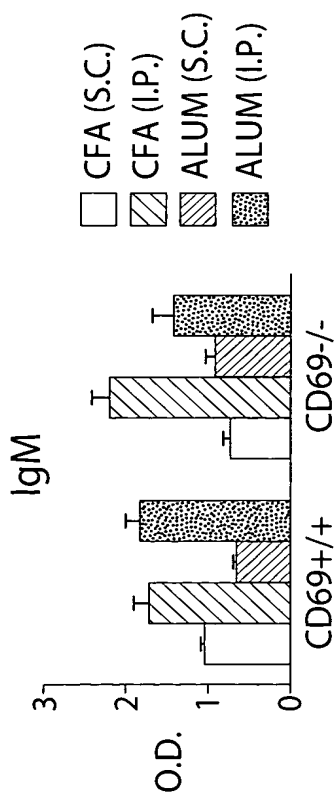


Fig. 18B

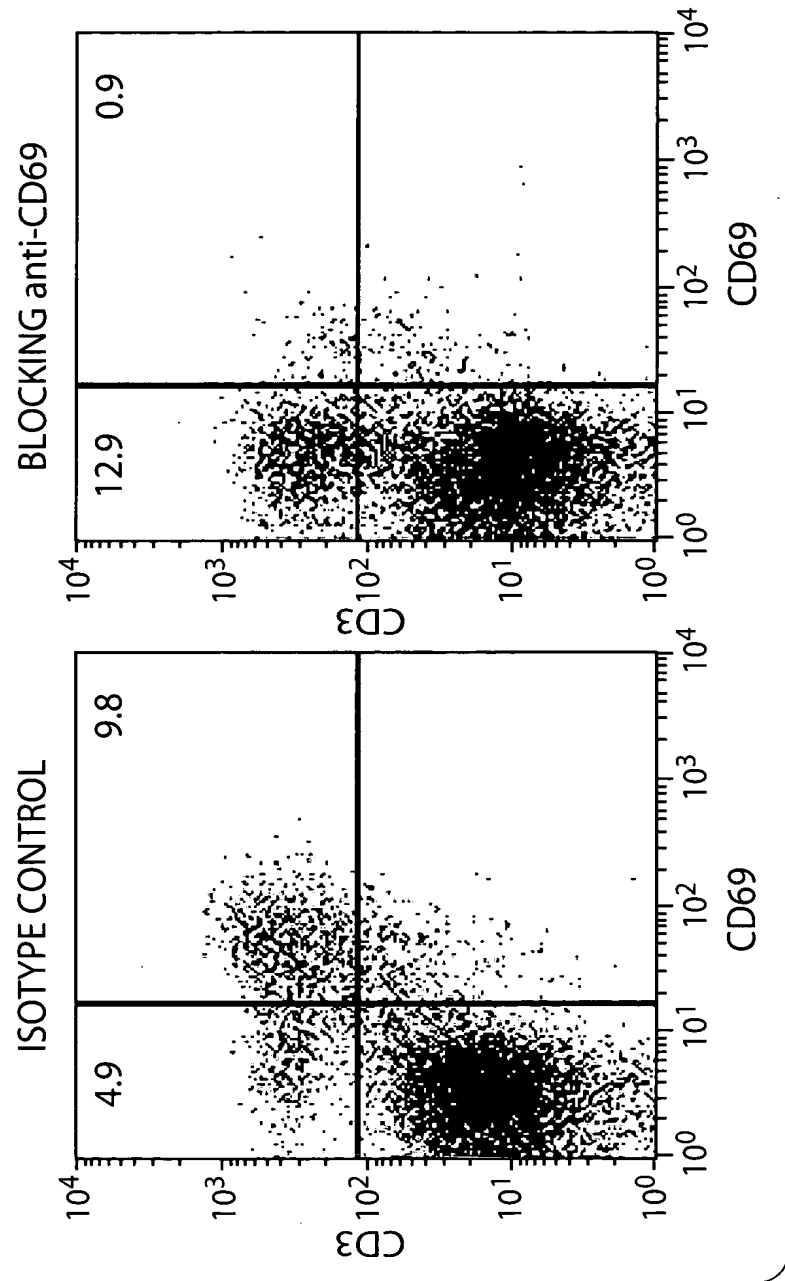


Fig. 19

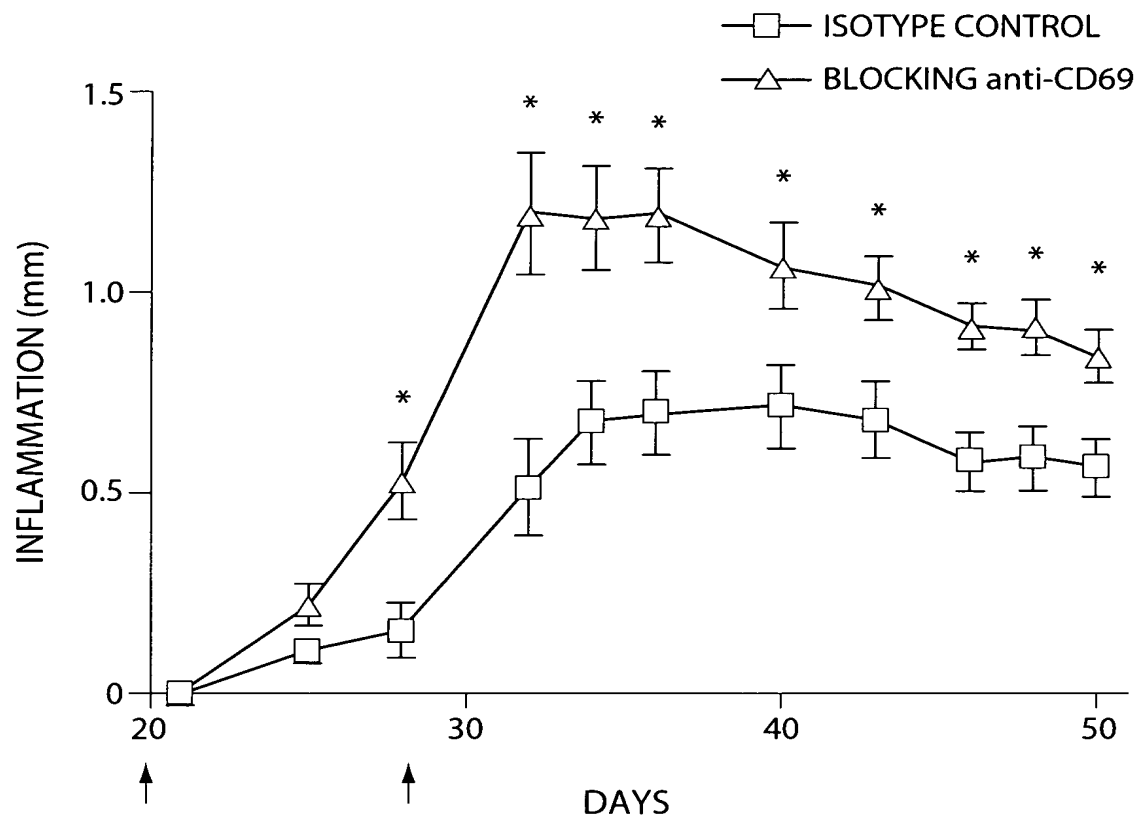


Fig. 20

CIA TREATMENT WITH DEPLETING anti-CD69

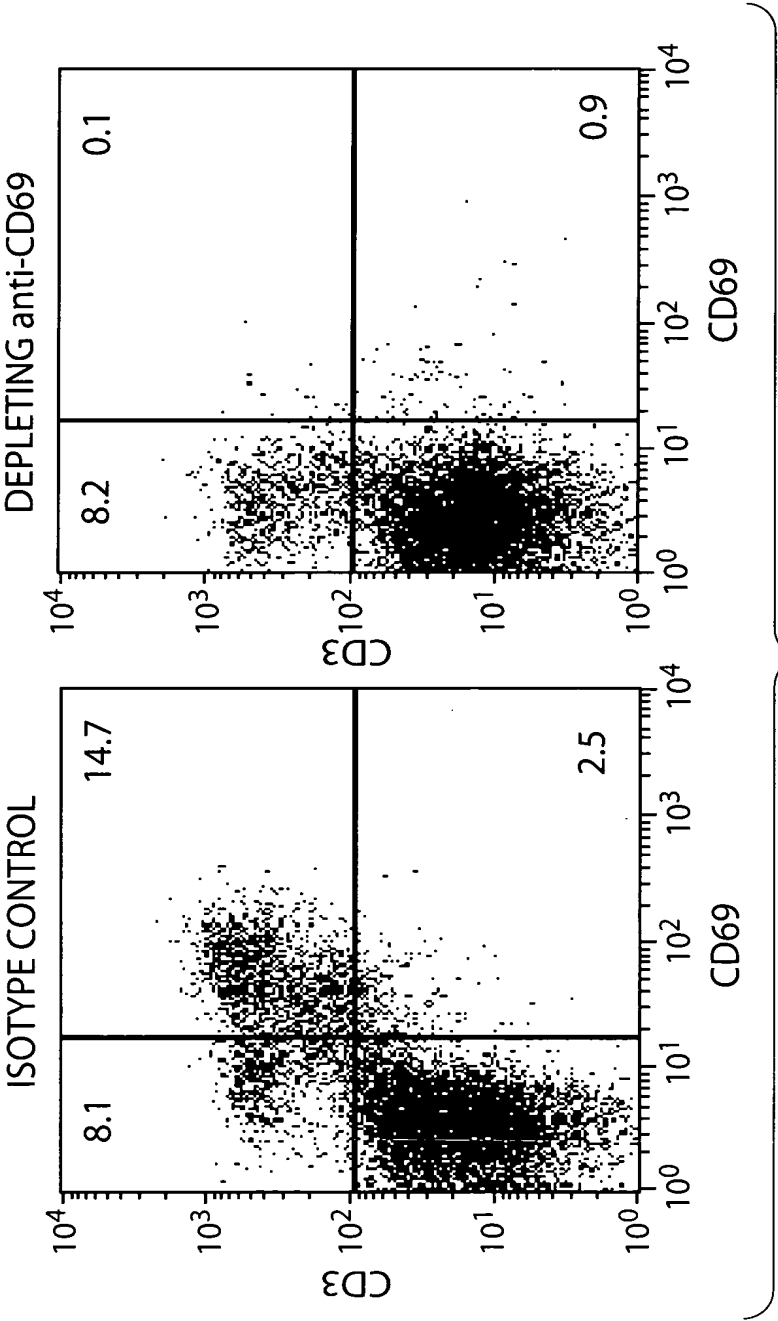


Fig. 21

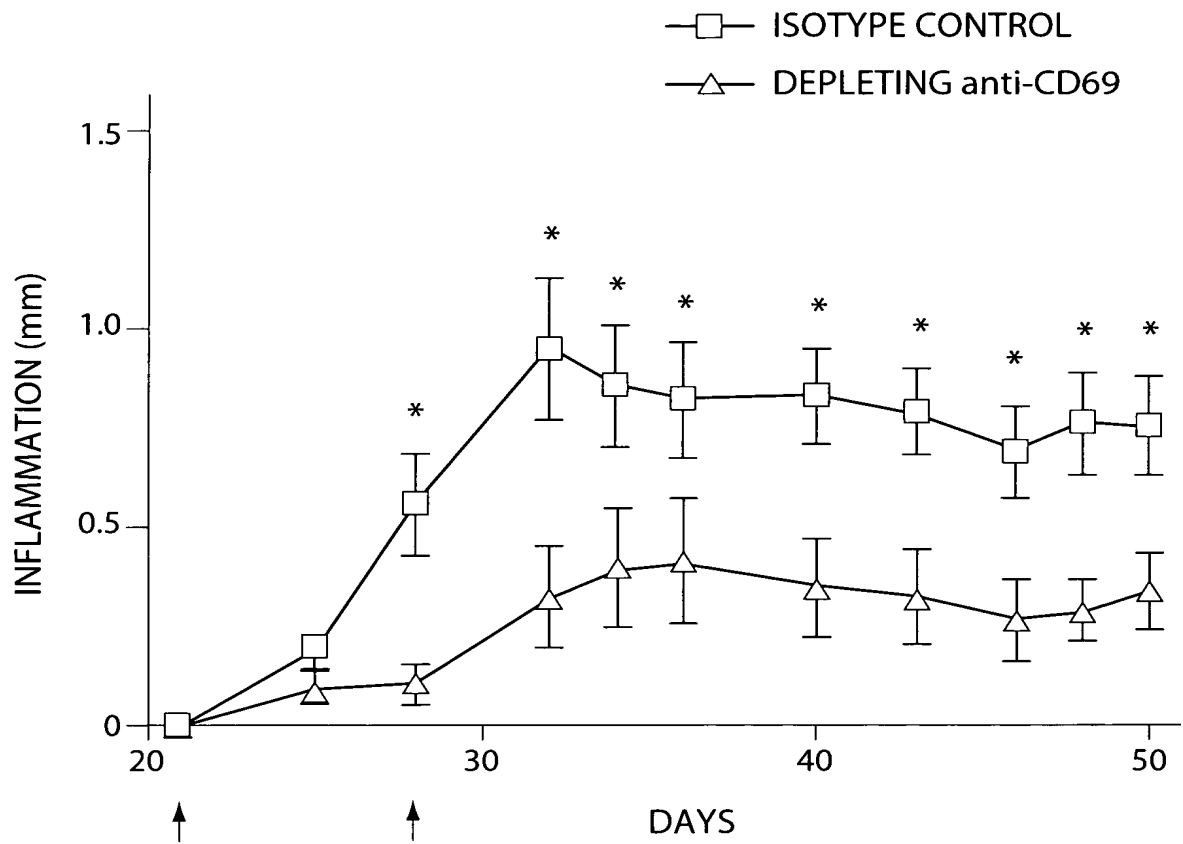


Fig. 22

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121 avidsekdmn flkryagree hwwglkkepg hpwkwsngke fnnwfnvtgs dkcvflknte
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Fig. 23

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301 aattcaagta aatacaactg ttccactcaa catgccgacc taactataat tgacaacata
361 gaagaaatga attttcttag gcggtataaa tgcagttctg atcactggat tggactgaag
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481 atgagaggga gtgaaggatg tgcctacctc agcgatgatg gtgcagcaac agctagatgt
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601 ggggaaaata gaaaataaca ttattaagtg taaaaccagc aaagtacttt ttttaattaaa
661 caaagttcga gttttgtacc tgtctgggta attctgctta cgtgtcaggc tacacataaa
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Fig. 24

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121 fllrykgpsd hwiglsreqg qpwkwingte wtrqfpilga gecaylndkg assarhyter
181 kwicksdih v      (SEQ ID NO:6)

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481 tatectggga gcaggagagt gtgcctattht gaatgacaaa ggtgccagta gtgccaggca
541 ctacacagag aggaagtgga tttgttccaa atcagatata catgtctaga tgttacagca
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661 agaattttatt tcttatacca acaggtatat gaaaatatgc tcaatatcac taataactgg
721 gaaaatacaa atcaaaatca tagtaaaata ttacctgttt tcatgggtgct aatattacct
781 gttctccac tgctaatagac ataccgaga atgagtaatt tataaataaa agagatttaa
841 ttgaaaaaaa      (SEQ ID NO:5)

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Fig. 25